

P. VANDEREGT

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RAW SEQUENCE LISTING
PATENT APPLICATION US/09/228,020

TECH CENTER 1600/2900
DATE: 11/07/2000
TIME: 02:58:37

INPUT SET: S36079.raw

This Raw Listing contains the General
Information Section and up to the first 5 pages.

SEQUENCE LISTING

ENTERED

(1) General Information:

(i) APPLICANT: ISNER, Jeffrey
ASAHARA, Takayuki

(ii) TITLE OF INVENTION: METHODS FOR REGULATING ANGIOGENESIS
(as amended)

(iii) NUMBER OF SEQUENCES: 2

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: DIKE, BRONSTEIN, ROBERTS & CUSMAN, LLP
(B) STREET: 130 Water Street
(C) CITY: Boston
(D) STATE: MA
(E) COUNTRY: USA
(F) ZIP: 02109

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: Diskette
(B) COMPUTER: IBM Compatible
(C) OPERATING SYSTEM: DOS
(D) SOFTWARE: FastSEQ for Windows Version 2.0

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER: 09/228,020
(B) FILING DATE:
(C) CLASSIFICATION:

(vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: 08/744,882
(B) FILING DATE:

(viii) ATTORNEY/AGENT INFORMATION:

(A) NAME: Resnick, David S
(B) REGISTRATION NUMBER: 34,235
(C) REFERENCE/DOCKET NUMBER: 46963

(ix) TELECOMMUNICATION INFORMATION:

(A) TELEPHONE: 617-523-3400
(B) TELEFAX: 617-523-6440
(C) TELEX:

RAW SEQUENCE LISTING
PATENT APPLICATION US/09/228,020DATE: 11/07/2000
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(2) INFORMATION FOR SEQ ID NO:1:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 30 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

AAGACATTTT CGGGCTCACG CTGCGCACCC

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(2) INFORMATION FOR SEQ ID NO:2:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 30 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

TGGGGTAGGC ACTTTAGTAG TTCTCCTAAC

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SEQUENCE VERIFICATION REPORT
PATENT APPLICATION US/09/228,020

DATE: 11/07/2000
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PAGE: 1

SEQUENCE MISSING ITEM REPORT
PATENT APPLICATION US/09/228,020

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< < THERE ARE NO ITEMS MISSING > >

PAGE: 1

SEQUENCE CORRECTION REPORT
PATENT APPLICATION US/09/228,020

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